## SEQUENCE LISTING

## (1) GENERAL INFORMATION:

- (i) APPLICANT: MOZES, Edna WAISMAN, Ari
- (ii) TITLE OF INVENTION: SYNTHETIC PEPTIDES AND PHARMACEUTICAL
  COMPOSITIONS COMPRISING THEM FOR THE TREATMENT
  OF SYSTEMIC LUPUS ERYTHEMATOSUS (SLE)
- (iii) NUMBER OF SEQUENCES: 10
- (iv) CORRESPONDENCE ADDRESS:
  - (A) ADDRESSEE: BROWDY AND NEIMARK
  - (B) STREET: 624 Ninth Street N.W., Ste. 300
  - (C) CITY: Washington
  - (D) STATE: D.C.
  - (E) COUNTRY: United States of America
  - (F) ZIP: 20001
- (v) COMPUTER READABLE FORM:
  - (A) MEDIUM TYPE: Floppy disk
  - (B) COMPUTER: IBM PC compatible
  - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
  - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
- (vi) CURRENT APPLICATION DATA:
  - (A) APPLICATION NUMBER: US 08/913,994
  - (B) FILING DATE: 29-SEP-1997
- (vii) PRIOR APPLICATION DATA:
  - (A) APPLICATION NUMBER: PCT/US96/04206
  - (B) FILING DATE: 27-MAR-1996
- (vi) PRIOR APPLICATION DATA:
  - (A) APPLICATION NUMBER: IL 113,159
  - (B) FILING DATE: 28-MAR-1995
- (viii) ATTORNEY/AGENT INFORMATION:
  - (A) NAME: BROWDY, Roger L.
  - (B) REGISTRATION NUMBER: 25,618
  - (C) REFERENCE/DOCKET NUMBER: MOZES=2
- (ix) TELECOMMUNICATION INFORMATION:
  - (A) TELEPHONE: (202) 628-5197
  - (B) TELEFAX: (202) 737-3528
- (2) INFORMATION FOR SEQ ID NO: 1:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 20 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single

- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
  - (D) OTHER INFORMATION:/note= Xaa in position 5 is Met, Ala or Val; Xaa in position 6 is Gln, Asp, Glu, or Arg; Xaa in position 7 is Trp or Ala; Xaa in position 8 is Val or Ser; and Xaa in position 9 is Lys, Glu or Ala.
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

Thr Gly Tyr Tyr Xaa Xaa Xaa Xaa Gln Ser Pro Glu Lys Ser Leu
1 10 15

Glu Trp Ile Gly 20

- (2) INFORMATION FOR SEQ ID NO: 2:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 20 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (ix) FEATURE:
    - (D) OTHER INFORMATION:/note= Xaa in position 9 is Thr, Val or Ala; Xaa in position 10 is Thr, Val or Ala; Xaa in position 11 is Tyr or Phe; Xaa in position 12 is Asn or Asp; Xaa in position 13 is Gln or Glu; Xaa in position 14 is Lys or Glu; and Xaa in position 15 is Phe or Tyr.
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

Glu Ile Asn Pro Ser Thr Gly Gly Xaa Xaa Xaa Xaa Xaa Xaa Xaa Lys 1 5 10 15

Ala Lys Ala Thr 20

- (2) INFORMATION FOR SEQ ID NO: 3:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 20 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (ix) FEATURE:

- (D) OTHER INFORMATION:/note= Xaa in position 6 is Phe, Thr or Gly; Xaa in position 7 is Leu, Ala or Ser; Xaa in position 8 is Trp or Ala; Xaa in position 9 is Glu or Lys; Xaa in position 13 is Met or Ala; and Xaa in position 14 is Asp, Lys or Ser.
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

Tyr Tyr Cys Ala Arg Xaa Xaa Xaa Pro Tyr Ala Xaa Xaa Tyr Trp

1 10 15

Gly Gln Gly Ser

- (2) INFORMATION FOR SEQ ID NO: 4:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 19 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (ix) FEATURE:
    - (D) OTHER INFORMATION:/note= Xaa in position 4 is Met or Ala; Xaa in position 5 is Asn, Asp or Arg; Xaa in position 6 is Trp or Ala; Xaa in position 7 is Val or Ser; Xaa in position 8 is Lys or Glu; Xaa in position 9 is Gln or Ala; Xaa in position 13 is Lys or Glu; and Xaa in position 14 is Ser or Ala.
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:

Gly Tyr Asn Xaa Xaa Xaa Xaa Xaa Ser His Gly Xaa Xaa Leu Glu
1 10 15

Trp Ile Gly

- (2) INFORMATION FOR SEQ ID NO: 5:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 18 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (ix) FEATURE:
    - (D) OTHER INFORMATION:/note= Xaa in position 6 is Ser or Phe; Xaa in position 7 is Gly or Ala; Xaa in position 8 is Arg, Ala or Glu; Xaa in position 11 is Asn or Asp; Xaa in position 12 is Tyr or Phe; and Xaa in position 13 is Trp, His or Ala.

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:

Tyr Tyr Cys Ala Arg Xaa Xaa Xaa Tyr Gly Xaa Xaa Xaa Gly Gln Gly
1 5 10 15

Thr Leu

- (2) INFORMATION FOR SEQ ID NO: 6:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 20 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:

Thr Gly Tyr Tyr Met Gln Trp Val Lys Gln Ser Pro Glu Lys Ser Leu 1 5 10 15

Glu Trp Ile Gly 20

- (2) INFORMATION FOR SEQ ID NO: 7:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 20 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:

Glu Ile Asn Pro Ser Thr Gly Gly Thr Thr Tyr Asn Gln Lys Phe Lys

1 10 15

Ala Lys Ala Thr 20

- (2) INFORMATION FOR SEQ ID NO: 8:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 20 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:

Tyr Tyr Cys Ala Arg Phe Leu Trp Glu Pro Tyr Ala Met Asp Tyr Trp 1 5 10 15

- (2) INFORMATION FOR SEQ ID NO: 9:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 19 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:

Gly Tyr Asn Met Asn Trp Val Lys Gln Ser His Gly Lys Ser Leu Glu 1 5 10 15

Trp Ile Gly

- (2) INFORMATION FOR SEQ ID NO: 10:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 18 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:

Tyr Tyr Cys Ala Arg Ser Gly Arg Tyr Gly Asn Tyr Trp Gly Gln Gly
1 5 10 15

Thr Leu